RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

10/541.657
PUT.
3/29/06

ENTERED



PCT

RAW SEQUENCE LISTING DATE: 03/29/2006
PATENT APPLICATION: US/10/541,657 TIME: 09:40:25

Input Set : A:\aren34.us5.pct sequence listing.ST25.txt
Output Set: N:\CRF4\03292006\J541657.raw

```
3 <110> APPLICANT: Jones, Robert M.
             Semple, Graeme
      5
             Fioravanti, Beatriz
             Pereira, Guilherme
      6
             Calderon, Imelda
      7
     8
             Uy, Jane
     9
             Duvvuri, Kameshwari
     10
             Choi, Jin Sun Karoline
     11
             Xiong, Yifeng
     12
              Vibha, Dave
     14 <120> TITLE OF INVENTION: 1,2,3-TRISUBSTITUTED ARYL AND HETEROARYL
DERIVATIVES AS
             MODULATORS OF METABOLISM AND THE PRPPHYLAXIS AND TREATMENT OF
     15
             DISORDERS RELATED THERETO SUCH AS DIABETES AND HYPERGLYCEMIA
     16
     18 <130> FILE REFERENCE: 34.US5.PCT
C--> 20 <140> CURRENT APPLICATION NUMBER: US/10/541,657
C--> 20 <141> CURRENT FILING DATE: 2005-07-07
     20 <150> PRIOR APPLICATION NUMBER: US 60/440394
     21 <151> PRIOR FILING DATE: 2003-01-14
     23 <150> PRIOR APPLICATION NUMBER: US 60/449,829
     24 <151> PRIOR FILING DATE: 2003-02-24
     26 <150> PRIOR APPLICATION NUMBER: US 60/453,390
     27 <151> PRIOR FILING DATE: 2003-03-06
     29 <150> PRIOR APPLICATION NUMBER: US 60/470,875
     30 <151> PRIOR FILING DATE: 2003-05-14
     32 <160> NUMBER OF SEO ID NOS: 6
     34 <170> SOFTWARE: PatentIn version 3.3
     36 <210> SEQ ID NO: 1
     37 <211> LENGTH: 1191
     38 <212> TYPE: DNA
     39 <213> ORGANISM: Homo Sapien
     41 <400> SEQUENCE: 1
     42 atgtacccat acgacgtccc agactacgct ggaagcttgg aatcatcttt ctcatttgga
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     44 gtgatccttg ctgtcctggc ctccctcatc attgctacta acacactagt ggctgtggct
                                                                              120
     46 gtgctgctgt tgatccacaa gaatgatggt gtcagtctct gcttcacctt gaatctggct
                                                                              180
     48 gtggctgaca ccttgattgg tgtggccatc tctggcctac tcacagacca gctctccagc
                                                                              240
                                                                              300
     50 ccttctcggc ccacacagaa gaccctgtgc agcctgcgga tggcatttgt cacttcctcc
                                                                              360
     52 qeagetgeet etgteeteae ggteatgetg ateacetttg acaggtacet tgeeateaag
                                                                              420
     54 caqceettee qetacttqaa qatcatqaqt qqqtteqtqq ceqqqqeetq cattqeeqqq
     56 ctqtqqttaq tqtcttacct cattggcttc ctcccactcg gaatccccat gttccagcag
                                                                              480
     58 actgcctaca aagggcagtg cagcttcttt gctgtatttc accctcactt cgtgctgacc
                                                                              540
                                                                              600
     60 ctctcctgcg ttggcttctt cccagccatg ctcctctttg tcttcttcta ctgcgacatg
                                                                              660
     62 ctcaagattg cctccatgca cagccagcag attcgaaaga tggaacatgc aggagccatg
     64 qctggaggtt atcgatcccc acggactccc agcgacttca aagctctccg tactgtgtct
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780
66 qttctcattq qqaqctttgc tctatcctgg acccccttcc ttatcactgg cattgtgcag
                                                                         840
68 gtggcctgcc aggagtgtca cctctaccta gtgctggaac ggtacctgtg gctgctcggc
70 gtgggcaact ccctgctcaa cccactcatc tatgcctatt ggcagaagga ggtgcgactg
                                                                         900
72 cagetetace acatggeect aggagtgaag aaggtgetea ceteatteet cetettete
                                                                         960
74 teggecagga attgtggece agagaggece agggaaagtt cetgteacat egteactate
                                                                        1020
76 tccagctcag agtttgatgg cgaattcgga tccaagggca attctgcaga tatccagcac
                                                                        1080
78 agtggcggcc gctcgagtct agagggcccg cggttcgaag gtaagcctat ccctaaccct
                                                                        1140
80 ctcctcggtc tcgattctac gcgtaccggt catcatcacc atcaccattg a
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83 <210> SEQ ID NO: 2
84 <211> LENGTH: 396
85 <212> TYPE: PRT
86 <213> ORGANISM: Homo Sapien
88 <400> SEQUENCE: 2
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91 1
                   5
94 Phe Ser Phe Gly Val Ile Leu Ala Val Leu Ala Ser Leu Ile Ile Ala
               20
                                   25
98 Thr Asn Thr Leu Val Ala Val Ala Val Leu Leu Ile His Lys Asn
                               40
102 Asp Gly Val Ser Leu Cys Phe Thr Leu Asn Leu Ala Val Ala Asp Thr
106 Leu Ile Gly Val Ala Ile Ser Gly Leu Leu Thr Asp Gln Leu Ser Ser
                        70
                                            75
110 Pro Ser Arg Pro Thr Gln Lys Thr Leu Cys Ser Leu Arg Met Ala Phe
                    85
                                        90
114 Val Thr Ser Ser Ala Ala Ala Ser Val Leu Thr Val Met Leu Ile Thr
                100
                                    105
                                                         110
118 Phe Asp Arg Tyr Leu Ala Ile Lys Gln Pro Phe Arg Tyr Leu Lys Ile
           115
                                120
122 Met .Ser Gly Phe Val Ala Gly Ala Cys Ile Ala Gly Leu Trp Leu Val
                            135
                                                140
126 Ser Tyr Leu Ile Gly Phe Leu Pro Leu Gly Ile Pro Met Phe Gln Gln
127 145
                        150
                                            155
130 Thr Ala Tyr Lys Gly Gln Cys Ser Phe Phe Ala Val Phe His Pro His
                                        170
                    165
134 Phe Val Leu Thr Leu Ser Cys Val Gly Phe Phe Pro Ala Met Leu Leu
                                    185
                180
138 Phe Val Phe Phe Tyr Cys Asp Met Leu Lys Ile Ala Ser Met His Ser
            195
                                200
                                                    205
142 Gln Gln Ile Arg Lys Met Glu His Ala Gly Ala Met Ala Gly Gly Tyr
                            215
                                             · 220
146 Arg Ser Pro Arg Thr Pro Ser Asp Phe Lys Ala Leu Arg Thr Val Ser
                                            235
150 Val Leu Ile Gly Ser Phe Ala Leu Ser Trp Thr Pro Phe Leu Ile Thr
                    245
                                        250
154 Gly Ile Val Gln Val Ala Cys Gln Glu Cys His Leu Tyr Leu Val Leu
                260
                                    265
158 Glu Arg Tyr Leu Trp Leu Leu Gly Val Gly Asn Ser Leu Leu Asn Pro
            275
                                280
```

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```
162 Leu Ile Tyr Ala Tyr Trp Gln Lys Glu Val Arg Leu Gln Leu Tyr His
                            295
       290
166 Met Ala Leu Gly Val Lys Lys Val Leu Thr Ser Phe Leu Leu Phe Leu
                                            315
170 Ser Ala Arg Asn Cys Gly Pro Glu Arg Pro Arg Glu Ser Ser Cys His
                                        330
                    325
174 Ile Val Thr Ile Ser Ser Glu Phe Asp Gly Glu Phe Gly Ser Lys
                340
                                    345
178 Gly Asn Ser Ala Asp Ile Gln His Ser Gly Gly Arg Ser Ser Leu Glu
            355
                                360
182 Gly Pro Arg Phe Glu Gly Lys Pro Ile Pro Asn Pro Leu Leu Gly Leu
     370
                            375
                                                380
186 Asp Ser Thr Arg Thr Gly His His His His His
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191 <211> LENGTH: 24
192 <212> TYPE: DNA
193 <213> ORGANISM: Artificial
195 <220> FEATURE:
196 <223> OTHER INFORMATION: Homo Sapien Primer
198 <400> SEQUENCE: 3
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199 cattgccggg ctgtggttag tgtc
202 <210> SEQ ID NO: 4
203 <211> LENGTH: 24
204 <212> TYPE: DNA
205 <213> ORGANISM: Artificial
207 <220> FEATURE:
208 <223> OTHER INFORMATION: Homo Sapien Primer
210 <400> SEQUENCE: 4
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211 ggcatagatg agtgggttga gcag
214 <210> SEQ ID NO: 5
215 <211> LENGTH: 22
216 <212> TYPE: DNA
217 <213> ORGANISM: Artificial
219 <220> FEATURE:
220 <223> OTHER INFORMATION: Rat Primer
222 <400> SEQUENCE: 5
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223 catgggccct gcaccttctt tg
226 <210> SEQ ID NO: 6
227 <211> LENGTH: 24
228 <212> TYPE: DNA
229 <213> ORGANISM: Artificial
231 <220> FEATURE:
232 <223> OTHER INFORMATION: Rat Primer
234 <400> SEQUENCE: 6
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235 gctccggatg gctgatgata gtga
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 03/29/2006 PATENT APPLICATION: US/10/541,657

TIME: 09:40:26

Input Set : A:\aren34.us5.pct sequence listing.ST25.txt

Output Set: N:\CRF4\03292006\J541657.raw

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:3,4,5,6

VERIFICATION SUMMARY

..

DATE: 03/29/2006

PATENT APPLICATION: US/10/541,657

TIME: 09:40:26

Input Set : A:\aren34.us5.pct sequence listing.ST25.txt

Output Set: N:\CRF4\03292006\J541657.raw

L:20 M:270 C: Current Application Number differs, Replaced Current Application No L:20 M:271 C: Current Filing Date differs, Replaced Current Filing Date